## SEQUENCE LISTING

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<110> Ausubel, Frederick M.
     Staskawicz, Brian J.
     Brent, Andrew F.
     Dahlbeck, Douglas
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     Kunkel, Barbara N.
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     Yu, Guo-Liang
<120> RPS2 GENE FAMILY, PRIMERS, PROBES, AND
 DETECTION METHODS
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<150> US 09/867,852
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<151> 2001-05-29

<150> US 09/301,085

<151> 1999-04-28

<150> US 08/310,912

<151> 1994-09-22

<150> US 08/227,360

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Ser Ala Ser Arg Trp Ser Thr Val Gly Phe Ile Leu Gly Arg Glu Gly
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Asp Arg Arg Lys Gln Ser Phe Glu Asp Ile Gln Ser Phe Glu Thr Glu
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Glu Glu Thr Arg Val Gly Ala Val Leu
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Gly Met Glu Lys Arg Ser Phe Arg Val Ile Ile Asn Ser Pro Ala Arg
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Gly Asp Tyr Ser Glu
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Met Trp Arg Ile Ala Thr Ser Val Asp His Phe Arg Arg Ser His Gly
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Ser
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Ile Gln Leu Arg Gln Pro Arg Glu
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Phe Tyr Arg Asp Arg Ala Ala Cys
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His His Leu Gln Gly Ile Phe Ser His Trp Gly Ser Glu Ser Gly Met
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Lys Thr His Met Pro Glu Thr Asp Asn Thr Asp Ala Pro Thr Glu Gly
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Leu Phe Glu Glu Asp Ser Asn Arg Val Phe His Ala Tyr Ala Cys Ser
Gln Ser Leu Gly Leu Val Val His Lys Tyr His
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Cys Gly Gln Lys Leu Cys Ile Val Asp Gly Ile
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Gly Ala Asp Pro Ser
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Gln Asn Pro Asp Leu Ala
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Asp Ser Val Val Tyr Gln Val Phe Gly Gly Val Val Ser Ser Val Tyr
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Val Arg Asn Lys Asp Lys Cys Ile Ala Thr Gly Ala Trp Glu Ser
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Lys Thr Glu Ala Ser Gly Pro Thr Lys Asn Ser Val Ser Ser Asp Asp
Pro Thr Arg Cys His Met Leu Ala Glu Gln Ala Arg Gly Ser Glu Leu
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                               25
Val Leu Gln Leu Arg Arg Leu Gly Thr Ala Glu Leu Trp Arg Arg
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Ser Arg Arg Thr Arg Ile Arg

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Ile Gly Asp Pro Lys Asn Ser Leu Arg Val Arg Cys Phe Ala
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Thr Tyr Thr Ala Ser Pro Arg
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Thr Pro Leu Leu Gln Ser Pro Ile Thr His
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Pro Trp Gln Glu Pro Glu Lys Thr
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Leu Gly-Val Pro Gly His Thr Arg Arg Phe
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Asn Gln Ser Val Gly Lys Phe Cys Lys Pro Arg Leu Ser Ala Glu Tyr
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Pro Leu His Lys His Phe Thr Leu Gln Gln Ala Glu Glu Cys Leu Met
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Gly Ser Glu Thr Pro Lys Ala Arg Gly Asp
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Thr Val Arg Leu Gln Arg Asp Arg Gly Ile Asp Lys Arg Thr Arg Glu
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Ser Ile Arg Arg Arg Ser Asn Ile Val Pro Lys Pro Glu Asp Leu Glu
Asn
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<212> PRT
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Gly Ser Ala Arg Thr Lys Gln His Pro Pro Ile Ser Ile Phe Ile Pro
1
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Lys Ser
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Asn Ile Ser His His Lys Leu Pro Gln Ser
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Glu Thr Ala Val Ser Gly Glu Glu Asp Pro Asp Glu Leu Ala Asn Ser

<210> 101

Leu Leu

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Thr Ser His His
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Glu Leu Arg Ala Leu Cys Thr Asn Met Ser Ile His Lys Met
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Gln Glu Ala Arg Lys Val Val Pro Val Lys Ser Ser Thr Phe His Ile
Ala Thr Lys Leu Glu Ile Met
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Lys Pro Asn Tyr Pro Arg
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gcgcatcatg gagctatttg gaattttcgc agggttatcg attcgtagtg ggaacccatt
                                                                        120
cattgtttgg aaccaccaac ggacgactta-acaagetccc-cgaggtgcat gatgaaaatt
                                                                        180
gctccagttg ccataaatca cagcccgctc agcagggagg tcccgtcaca cgcggcaccc
                                                                        240
                                                                        300
actcaggcaa agcaaaccaa ccttcaatct gaagctggcg atttagatgc aagaaaaagt
agegetteaa geeeggaaac eegggeatta etegetaeta agacagtaet egggagacae
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aagatagagg ttccggcctt tggagggtgg ttcaaaaaga aatcatctaa gcacgagacg
                                                                        420
                                                                        480
ggcggttcaa gtgccaacgc agatagttcg agcgtggctt ccgattccac cgaaaaacct
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ttgttccgtc tcacgcacgt tccttacgta tcccaaggta atgagcgaat gggatgttgg
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tatgcctgcg caagaatggt tggccattct gtcgaagctg ggcctcgcct agggctgccg
                                                                        660
gagctctatg agggaaggga ggcgccagct gggctacaag atttttcaga tgtagaaagg
                                                                        720
tttattcaca atgaaggatt aactegggta gacettecag acaatgagag atttacacae
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gaagagttgg gtgcactgtt gtataagcac gggccgatta tatttgggtg gaaaactccg
                                                                     780
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aatgacaget ggcacatgte ggteeteact ggtgtegata aagagaegte gteeattact
tttcacqatc cccgacaqqq qccqqaccta qcaatqccgc tcgattactt taatcagcqa
                                                                     900
ttggcatggc aggttccaca cgcaatgctc taccgctaag tagcagggta tcttcacgtg
                                                                     960
geggeateat gaeaageeea tgatgeegee ageagetace tgaatgeegt etggettttt
                                                                    1020
ggtccctatt gtcgtatccg gaagatgacg tcaaagaatc tcggcaagag ctttcttgct
                                                                    1080
cqactectea getteeggat egateaggte gettgecaga gegegettgt ceatgageat
                                                                    1140
ctgccacage tgctggtcga tggtgtcctc agctaaaggg attttgacga caaccatgcg
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caactgcccg ttgcgatacg ctcgatcctg aagccccggt gtccatggca gccccaagaa
                                                                    1260
aaagacatag ttcgccgctg tgaggttgta gcctgtgccg gcggccgacc tggtcccgat
                                                                    1320
aaacaccctg cagtccggat cctgctggaa agcatcaatc gccttctgcc gcttcttggg
                                                                    1380
cqaqtcactg cccaccaacg tcacqcaccc gacqccaaqc ttqagqcaqt gctcccqcaa
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                                                                    1491
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                           40
Glu Thr Arg Ala Leu Leu Ala Thr Lys Thr Val Leu Gly Arg His Lys
Ile Glu Val Pro Ala Phe Gly Gly Trp Phe Lys Lys Lys Ser Ser Lys
His Glu Thr Gly Gly Ser Ser Ala Asn Ala Asp Ser Ser Ser Val Ala
Ser Asp Ser Thr Glu Lys Pro Leu Phe Arg Leu Thr His Val Pro Tyr
                               105
Val Ser Gln Gly Asn Glu Arg Met Gly Cys Trp Tyr Ala Cys Ala Arg
                           120
Met Val Gly His Ser Val Glu Ala Gly Pro Arg Leu Gly Leu Pro Glu
                       135
Leu Tyr Glu Gly Arg Glu Ala Pro Ala Gly Leu Gln Asp Phe Ser Asp
                    150
                                       155
Val Glu Arg Phe Ile His Asn Glu Gly Leu Thr Arg Val Asp Leu Pro
                                   170
Asp Asn Glu Arg Phe Thr His Glu Glu Leu Gly Ala Leu Leu Tyr Lys
            180
                               185
His Gly Pro Ile Ile Phe Gly Trp Lys Thr Pro Asn Asp Ser Trp His
                           200
                                               205
Met Ser Val Leu Thr Gly Val Asp Lys Glu Thr Ser Ser Ile Thr Phe
                       215
                                           220
His Asp Pro Arg Gln Gly Pro Asp Leu Ala Met Pro Leu Asp Tyr Phe
Asn Gln Arg Leu Ala Trp Gln Val Pro His Ala Met Leu Tyr Arg
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<sup>&</sup>lt;211> 1258

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Arabidopsis thaliana

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                            40
Ser Thr Asn Pro Ser Gly Ser Phe Pro Ser Val Glu Tyr Glu Val Phe
Leu Ser Phe Arg Gly Pro Asp Thr Arg Glu Gln Phe Thr Asp Phe Leu
Tyr Gln Ser Leu Arg Arg Tyr Lys Ile His Thr Phe Arg Asp Asp Asp
Glu Leu Leu Lys Gly Lys Glu Ile Gly Pro Asn Leu Leu Arg Ala Ile
Aşp Gln Ser Lyş Ile Tyr Val Pro Ile Ile Ser Ser Gly Tyr Ala Aşp
                            120
Ser Lys Trp Cys Leu Met Glu Leu Ala Glu Ile Val Arg Arg Gln Glu
                        135
Glu Asp Pro Arg Arg Ile Ile Leu Pro Ile Phe Tyr Met Val Asp Pro
                    150
                                       155
Ser Asp Val Arg His Gln Thr Gly Cys Tyr Lys Lys Ala Phe Arg Lys
               - 165
                                    170
His Ala Asn Lys Phe Asp Gly Gln Thr Ile Gln Asn Trp Lys Asp Ala
            180
                                185
Leu Lys Lys Val Gly Asp Leu Lys Gly Trp His Ile Gly Lys Asn Asp
                            200
Lys Gln Gly Ala Ile Ala Asp Lys Val Ser Ala Asp Ile Trp Ser His
                        215
Ile Ser Lys Glu Asn Leu Ile Leu Glu Thr Asp Glu Leu Val Gly Ile
                    230
                                        235
Asp Asp His Ile Thr Ala Val Leu Glu Lys Leu Ser Leu Asp Ser Glu
Asn Val Thr Met Val Gly Leu Tyr Gly Met Gly Gly Ile Gly Lys Thr
Thr Thr Ala Lys Ala Val Tyr Asn Lys Ile Ser Ser Cys Phe Asp Cys
                            280
Cys Cys Phe Ile Asp Asn Ile Arg Glu Thr Gln Glu Lys Asp Gly Val
                        295
Val Val Leu Gln Lys Lys Leu Val Ser Glu Ile Leu Arg Ile Asp Ser
                                        315
Gly Ser Val Gly Phe Asn Asn Asp Ser Gly Gly Arg Lys Thr Ile Lys
                                    330
Glu Arg Val Ser Arg Phe Lys Ile Leu Val Val Leu Asp Asp Val Asp
            340
                                345
Glu Lys Phe Lys Phe Glu Asp Met Leu Gly Ser Pro Lys Asp Phe Ile
                                                365
                            360
Ser Gln Ser Arg Phe Ile Ile Thr Ser Arg Ser Met Arg Val Leu Gly
                        375
                                            380
Thr-Leu-Asn-Glu-Asn-Gln-Cys-Lys-Leu-Tyr-Glu-Val.Gly-Ser-Met_Ser_
                    390
                                        395
Lys Pro Arg Ser Leu Glu Leu Phe Ser Lys His Ala Phe Lys Lys Asn
                                    410
Thr Pro Pro Ser Ser Tyr Tyr Glu Thr Leu Ala Asn Asp Val Val Asp
                                425
Thr Thr Ala Gly Leu Pro Leu Thr Leu Lys Val Ile Gly Ser Leu Leu
                            440
Phe Lys Gln Glu Ile Ala Val Trp Glu Asp Thr Leu Glu Gln Leu Arg
    450
                        455
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Arg Thr Leu Asn Leu Asp Glu Val Tyr Asp Arg Leu Lys Ile Ser Tyr
                   470
Asp Ala Leu Asn Pro Glu Ala Lys Glu Ile Phe Leu Asp Ile Ala Cys
                                   490
               485
Phe Phe Ile Gly Gln Asn Lys Glu Glu Pro Tyr Tyr Met Trp Thr Asp
                               505
           500
Cys Asn Phe Tyr Pro Ala Ser Asn Ile Ile Phe Leu Ile Gln Arg Cys
                           520
Met Ile Gln Val Gly Asp Asp Asp Glu Phe Lys Met His Asp Gln Leu
                       535
Arg Asp Met Gly Arg Glu Ile Val Arg Arg Glu Asp Val Leu Pro Trp
                   550
Lys Ser Arg Ile Trp Ser Ala Glu Glu Gly Ile Asp Leu Leu Asn
                                   570
               565
Lys Arg Lys Gly Ser Ser Lys Val Lys Ala Ile Ser Ile Pro Trp Gly
                               585
Val Lys Tyr Glu Phe Lys Ser Glu Cys Phe Leu Asn Leu Ser Glu Leu
                           600
Arg Tyr Leu His Ala Arg Glu Ala Met Leu Thr Gly Asp Phe Asn Asn
                                           620
                       615
Leu Leu Pro Asn Leu Lys Trp Leu Glu Leu Pro Phe Tyr Lys His Gly
                                        635
                   630
Glu Asp Asp Pro Pro Leu Thr Asn Tyr Thr Met Lys Asn Leu Ile Ile
                                    650
               645
Val Ile Leu Glu His Ser His Ile Thr Ala Asp Asp Trp Gly Gly Trp
                                665
            660
Arg His Met Met Lys Met Ala Glu Arg Leu Lys Val Val Arg Leu Ala
                            680
Ser Asn Tyr Ser Leu Tyr Gly Arg Arg Val Arg Leu Ser Asp Cys Trp
                                            700
                        695
Arg Phe Pro Lys Ser Ile Glu Val Leu Ser Met Thr Ala Ile Glu Met
                    710
Asp Glu Val Asp Ile Gly Glu Leu Lys Lys Leu Lys Thr Leu Val Leu
                                   730
                725
Lys Pro Cys Pro Ile Gln Lys Ile Ser Gly Gly Thr Phe Gly Met Leu
                                745
Lys Gly Leu Arg Glu Leu Cys Leu Glu Phe Asn Trp Gly Thr Asn Leu
                          . 760
Arg Glu Val Val Ala Asp Ile Gly Gln Leu Ser Ser Leu Lys Val Leu
                        775
Lys Thr Gly Ala Lys Glu Val Glu Ile Asn Glu Phe Pro Leu Gly Leu
                    790
                                        795
Lys Thr Glu Leu Ser Thr Ser Ser Arg Ile Pro Asn Asn Leu Ser Gln
                805
                                    810
Leu Leu Asp Leu Glu Val Leu Lys Val Tyr Asp Cys Lys Asp Gly Phe
                                825
                                                   830 .
Asp Met Pro Pro Ala Ser Pro Ser Glu Asp Glu Ser Ser Val Trp Trp
                            840
                                                845
Lys Val Ser Lys Leu Lys Ser Leu Gln Leu Glu Lys Thr Arg Ile Asn
 850 ..... 860 .....
Val Asn Val Val Asp Asp Ala Ser Ser Gly Gly His Leu Pro Arg Tyr
                                        875
                    870
Leu Leu Pro Thr Ser Leu Thr Tyr Leu Lys Ile Tyr Gln Cys Thr Glu
                                    890
                885
Pro Thr Trp Leu Pro Gly Ile Glu Asn Leu Glu Asn Leu Thr Ser Leu
                                905
            900
Glu Val Asn Asp Ile Phe Gln Thr Leu Gly Gly Asp Leu Asp Gly Leu
                            920 -
Gln Gly Leu Arg Ser Leu Glu Ile Leu Arg Ile Arg Lys Val Asn Gly
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935
Leu Ala Arg Ile Lys Gly Leu Lys Asp Leu Leu Cys Ser Ser Thr Cys
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                  950
Lys Leu Arg Lys Phe Tyr Ile Thr Glu Cys Pro Asp Leu Ile Glu Leu
                                  970
               965
Leu Pro Cys Glu Leu Gly Val Gln Thr Val Val Val Pro Ser Met Ala
                              985
Glu Leu Thr Ile Arg Asp Cys Pro Arg Leu Glu Val Gly Pro Met Ile
                          1000
                                            1005
Arg Ser Leu Pro Lys Phe Pro Met Leu Lys Lys Leu Asp Leu Ala Val
                                        1020
                      1015
Ala Asn Ile Thr Lys Glu Glu Asp Leu Asp Ala Ile Gly Ser Leu Glu
                  1030
                                     1035
Glu Leu Val Ser Leu Glu Leu Glu Leu Asp Asp Thr Ser Ser Gly Ile
           1045
                                 1050
Glu Arg Ile Val Ser Ser Ser Lys Leu Gln Lys Leu Thr Thr Leu Val
                             1065
          1060
Val Lys Val Pro Ser Leu Arg Glu Ile Glu Gly Leu Glu Leu Lys
      1075
                          1080
Ser Leu Gln Asp Leu Tyr Leu Glu Gly Cys Thr Ser Leu Gly Arg Leu
           1095
                                         1100
Pro Leu Glu Lys Leu Lys Glu Leu Asp Ile Gly Gly Cys Pro Asp Leu
                 1110
                                     1115
Thr Glu Leu Val Gln Thr Val Val Ala Val Pro Ser Leu Arg Gly Leu
                                 1130
                                                     1135
              1125
Thr Ile Arg Asp Cys Pro Arg Leu Glu Val Gly Pro Met Ile Gln Ser
                1145
                                                 1150
          1140
Leu Pro Lys Phe Pro Met Leu Asn Glu Leu Thr Leu Ser Met Val Asn
                         1160
                                             1165
       1155
Ile Thr Lys Glu Asp Glu Leu Glu Val Leu Gly Ser Leu Glu Glu Leu
                      1175
                                          1180
Asp Ser Leu Glu Leu Thr Leu Asp Asp Thr Cys Ser Ser Ile Glu Arg
                                      1195
                   1190
Ile Ser Phe Leu Ser Lys Leu Gln Lys Leu Thr Thr Leu Ile Val Glu
                                  1210
               1205
Val Pro Ser Leu Arg Glu Ile Glu Gly Leu Ala Glu Leu Lys Ser Leu
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Arg Ile Leu Tyr Leu Glu Gly Cys Thr Ser Leu Glu Arg Leu Trp Pro
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Asp Gln Gln Leu Gly Ser Leu Lys Asn
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<211> 1143
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<212> PRT

<213> Arabidopsis thaliana

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Arg Trp Cys Leu Asn Glu Leu Val Lys Ile Met Glu Cys Lys Thr Arg
Phe Lys Gln Thr Val Ile Pro Ile Phe Tyr Asp Val Asp Pro Ser His
           100
                               105
Val Arg Asn Gln Lys Glu Ser Phe Ala Lys Ala Phe Glu Glu His Glu
                           120
Thr Lys Tyr Lys Asp Asp Val Glu Gly Ile Gln Arg Trp Arg Ile Ala
                       135
Leu Asn Glu Ala Ala Asn Leu Lys Gly Ser Cys Asp Asn Arg Asp Lys
                   150
Thr Asp Ala Asp Cys Ile Arg Gln Ile Val Asp Gln Ile Ser Ser Lys
                                   170
               165
Leu Cys Lys Ile Ser Leu Ser Tyr Leu Gln Asn Ile Val Gly Ile Asp
           180
                               185
Thr His Leu Glu Lys Ile Glu Ser Leu Leu Glu Ile Gly Ile Asn Gly
                           200
Val Arg Ile Met Gly Ile Trp Gly Met Gly Gly Val Gly Lys Thr Thr
                       215
Ile Ala Arg Ala Ile Phe Asp Thr Leu Leu Gly Arg Met Asp Ser Ser
                   230
                                      235
Tyr Gln Phe Asp Gly Ala Cys Phe Leu Lys Asp Ile Lys Glu Asn Lys
                                   250
               245
Arg Gly Met His Ser Leu Gln Asn Ala Leu Leu Ser Glu Leu Leu Arg
           260
                               265
Glu Lys Ala Asn Tyr Asn Asn Glu Glu Asp Gly Lys His Gln Met Ala
                           280
Ser Arg Leu Arg Ser Lys Lys Val Leu Ile Val Leu Asp Asp Ile Asp
                       295
Asn Lys Asp His Tyr Leu Glu Tyr Leu Ala Gly Asp Leu Asp Trp Phe
                   310
Gly Asn Gly Ser Arg Ile Ile Ile Thr Thr Arg Asp Lys His Leu Ile
               325
                                   330
Glu Lys Asn Asp Ile Ile Tyr Glu Val Thr Ala Leu Pro Asp His Glu
           340
                               345
Ser Ile Gln Leu Phe Lys Gln His Ala Phe Gly Lys Glu Val Pro Asn
                           360
Glu Asn Phe Glu Lys Leu Ser Leu Glu Val Val Asn Tyr Ala Lys Gly
                       37.5
                                           380
Leu Pro Leu Ala Leu Lys Val Trp Gly Ser Leu Leu His Asn Leu Arg
                   390
                                       395
Leu Thr Glu Trp Lys Ser Ala Ile Glu His Met Lys Asn Asn Ser Tyr
               405
                                   410
Ser Gly Ile Ile Asp Lys Leu Lys Ile Ser Tyr Asp Gly Leu Glu Pro
           420
                               425
Lys Gln Gln Glu Met Phe Leu Asp Ile Ala Cys Phe Leu Arg Gly Glu
       435
                           440
                                               445
Glu Lys Asp Tyr Ile Leu Gln Ile Leu Glu Ser Cys His Ile Gly Ala
                       455
                                           460
Glu Tyr Gly Leu Arg Ile Leu Ile Asp Lys Ser Leu Val Phe Ile Ser
Glu Tyr Asn Gln Val Gln Met His Asp Leu Ile Gln Asp Met Gly Lys
               485
                                   490
Tyr Ile Val Asn Phe Gln Lys Asp Pro Gly Glu Arg Ser Arg Leu Trp
                               505
Leu Ala Lys Glu Val Glu Glu Val Met Ser Asn Asn Thr Gly Thr Met
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Ala Met Glu Ala Ile Trp Val Ser Ser Tyr Ser Ser Thr Leu Arg Phe
                       535
Ser Asn Gln Ala Val Lys Asn Met Lys Arg Leu Arg Val Phe Asn Met
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                    550
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Gly Arg Ser Ser Thr His Tyr Ala Ile Asp Tyr Leu Pro Asn Asn Leu
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                                    570
Arg Cys Phe Val Cys Thr Asn Tyr Pro Trp Glu Ser Phe Pro Ser Thr
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Phe Glu Leu Lys Met Leu Val His Leu Gln Leu Arg His Asn Ser Leu
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Arg His Leu Trp Thr Glu Thr Lys His Leu Pro Ser Leu Arg Arg Ile
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Asp Leu Ser Trp Ser Lys Arg Leu Thr Arg Thr Pro Asp Phe Thr Gly
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Met Pro Asn Leu Glu Tyr Val Asn Leu Tyr Gln Cys Ser Asn Leu Glu
Glu Val His His Ser Leu Gly Cys Cys Ser Lys Val Ile Gly Leu Tyr
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Leu Asn Asp Cys Lys Ser Leu Lys Arg Phe Pro Cys Val Asn Val Glu
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Ser Leu Glu Tyr Leu Gly Leu Arg Ser Cys Asp Ser Leu Glu Lys Leu
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                        695
Pro Glu Ile Tyr Gly Arg Met Lys Pro Glu Ile Gln Ile His Met Gln
                    710
                                        715
Gly Ser Gly Ile Arg Glu Leu Pro Ser Ser Ile Phe Gln Tyr Lys Thr
                                    730
                725
His Val Thr Lys Leu Leu Trp Asn Met Lys Asn Leu Val Ala Leu
            740
                                745
Pro Ser Ser Ile Cys Arg Leu Lys Ser Leu Val Ser Leu Ser Val Ser
                           760
Gly Çys Ser Lys Leu Glu Ser Leu Pro Glu Glu Ile Gly Asp Leu Asp
                    775
                                            780
Asn Leu Arg Val Phe Asp Ala Ser Asp Thr Leu Ile Leu Arg Pro Pro
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                                        795
Ser Ser Ile Ile Arg Leu Asn Lys Leu Ile Ile Leu Met Phe Arg Gly
                                    810
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Phe Lys Asp Gly Val His Phe Glu Phe Pro Pro Val Ala Glu Gly Leu
                                825
His Ser Leu Glu Tyr Leu Asn Leu Ser Tyr Cys Asn Leu Ile Asp Gly
                            840
Gly Leu Pro Glu Glu Ile Gly Ser Leu Ser Ser Leu Lys Lys Leu Asp
                        855
Leu Ser Arg Asn Asn Phe Glu His Leu Pro Ser Ser Ile Ala Gln Leu
                    870
                                        875
Gly Ala Leu Gln Ser Leu Asp Leu Lys Asp Cys Gln Arg Leu Thr Gln
                                    890
Leu Pro Glu Leu Pro Pro Glu Leu Asn Glu Leu His Val Asp Cys His
                                905
Met Ala Leu Lys Phe Ile His Tyr Leu Val Thr Lys Arg Lys Lys Leu
                            920
His Arg Val Lys Leu Asp Asp Ala His Asn Asp Thr Met Tyr Asn Leu
                        935
                                            940
Phe Ala-Tyr-Thr Met Phe Gln Asn Ile Ser Ser Met Arg His Asp Ile
                    950
                                        955
Ser Ala Ser Asp Ser Leu Ser Leu Thr Val Phe Thr Gly Gln Pro Tyr
                965
                                    970
Pro Glu Lys Ile Pro Ser Trp Phe His His Gln Gly Trp Asp Ser Ser
                                985
Val Ser Val Asn Leu Pro Glu Asn Trp Tyr Ile Pro Asp Lys Phe Leu
                            1000
                                                1005
Gly Phe Ala Val Cys Tyr Ser Arg Ser Leu Ile Asp Thr Thr Ala His
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Pro Glu Val Asn Ala Leu Leu Gln Met Arg Glu Asn Ser Asn Glu Pro
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Lys Lys Lys Thr Leu Asp Ser Trp Lys Val Val Glu Gln Ser Leu Ser
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Ser Gln Arg Ile Gly Ser Leu Glu Glu Ser Ile Ser Ile Ile Gly Phe
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Ser Tyr Lys Asn Leu Pro His Tyr Leu Lys Pro Cys Phe Leu Tyr Phe
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Leu Ile Trp Val Gln Met Ser Arg Glu Phe Gly Glu Cys Thr Ile Gln
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Gln Ala Val Gly Ala Arg Leu Gly Leu Ser Trp Asp Glu Lys Glu Thr
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Gly Glu Asn Arg Ala Leu Lys Ile Tyr Arg Ala Leu Arg Gln Lys Arg
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Thr Gly Val Pro Arg Pro Asp Arg Glu Asn Lys Cys Lys Val Met Phe
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Ser Lys Val Trp Arg Lys Asp Leu Leu Glu Ser Ser Ile Arg Arg
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Ile Thr Leu Gly Gly Ala Met Ala His Arg Glu Thr Glu Glu Glu Trp
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Thr Val Leu Ser Leu Glu Thr Leu Lys Thr Leu Phe Glu Phe Gly Ala
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                                                                      1080
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gaattaaagg ataaactact tggaggatca cctgagcttg atgtcatctc aatcgttggc
                                                                      1320
atgccaggat tgggcaagac tacactagca aagaagattt acaatgatcc agaagtcacc
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                                                                      1620
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                                                                      1860
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                                                                      2880
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                                                                      4860
                                                                      4920
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tgaaccaaaa gttacctttt ttttttcttc ttaatggcat tactttgaag cacatgtttg
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                                                                           . 21
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 <213> Arabidopsis thaliana
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 narnggnarn cc
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 <213> Arabidopsis thaliana
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                                                                             17
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 ggnytnccny tndsnbt
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                                                                              2.6
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narnswytyn arytt
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                                                                    17
ggnytnmrnw snytnga
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Leu Lys Phe Ser Tyr Asp Asn Leu Glu Ser Asp Leu Leu
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<211> 16
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Gly Val Tyr Gly Pro Gly Gly Val Gly Lys Thr Thr Leu Met Gln Ser
                                     10
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<211> 14
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Gly Gly Leu Pro Leu Ala Leu Ile Thr Leu Gly Gly Ala Met
                5
<210> 191
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<212> PRT
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21

<213> Arabidopsis thaliana

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<221> VARIANT
<222> (2)...(2)
<223> Xaa is Met or Pro
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<222> (3)...(3)
<223> Xaa is Gly or Pro
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<222> (5)...(5)
<223> Xaa is Ile, Leu or Val
<221> VARIANT
<222> (10) ... (10)
<223> Xaa is Ile, Leu or Thr
<221> VARIANT
<222> (11)...(11)
<223> Xaa is Ala or Met
<400> 191
Gly Xaa Xaa Gly Xaa Gly Lys Thr Thr Xaa Xaa
<210> 192
<211> 11
<212> PRT
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<221> VARIANT
<222> (1)...(11)
<223> Xaa at 1 is Phe or Lys; Xaa at 2 is Arg or Lys;
      Xaa at 3 is Ile, Val or Phe; Xaa at 5 is Ile, Leu or Val; Xaa at 6 is Ile or Leu; Xaa at 7 is Ile or
       Val; Xaa at 10 is Ile, Leu or Val; Xaa at 11 is
       Asp or Trp;
<400> 192
Xaa Xaa Xaa Leu Xaa Xaa Xaa Asp Asp Xaa Xaa
                                       10
 1
<210> 193
<211> 8
<212> PRT
<213> Arabidopsis thaliana
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<221> VARIANT
<222> (1)...(8)
<223> Xaa at 1 is Ser or Cys; Xaa at 2 is Arg or Lys;
       Xaa at 3 is Phe, Ile or Val; Xaa at 4 is Ile or
       Met; Xaa at 5 is Ile, Leu or Phe; Xaa at 7 is Ser,
       Cys or Thr;
 <400> 193
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Xaa Xaa Xaa Xaa Thr Xaa Arg
1
<210> 194
<211> 8
<212> PRT
<213> Arabidopsis thaliana
<220>
<221> VARIANT
<222> (1) ... (8)
<223> Xaa at 5 is Thr, Ala or Thr; Xaa at 6 is Leu or
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      or Thr;
<400> 194
Gly Leu Pro Leu Xaa Xaa Xaa Xaa
<210> 195
<211> 7
<212> PRT
<213> Arabidopsis thaliana
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<221> VARIANT
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<223> Xaa at 1 is Lys or Gly; Xaa at 2 is Ile or Phe; Xaa at 5 is Asp or Lys; Xaa at 6 is Ala, Gly or
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Xaa Xaa Ser Tyr Xaa Xaa Leu
1
<210> 196 ·
<211> 4
<212> PRT
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Asn Ser His Arg
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<400> 197
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<210> 198
<211> 4
<212> PRT
<213> Arabidopsis thaliana
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Thr Gly Asp Leu
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<211> 4
<212> PRT
<213> Arabidopsis thaliana
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His Gly Thr Tyr
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<211> 11
<212> PRT
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Arg Met Ser His Gly Phe Arg Asn Ser Gln Ser
 1 5
<210> 201
<211> 27
<212> PRT
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<223> Xaa at position 5 is Met or Asp
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Pro Ile Phe Tyr Xaa Val Asp Pro Ser
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<211> 6
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<222> (1)...(6)
<223> Xaa at position 5 is Asp or Thr
<400> 203
Val Gly Ile Asp Xaa His
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<211> 9
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<221> VARIANT
<222> (1)...(9)
<223> Xaa at position 1 is Gln or Leu; Xaa at position 2
      is Leu or Ile; Xaa at position 3 is Arg or Gln.
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Met His Asp Xaa Xaa Xaa Asp Met Gly
                . 5
<210> 205
<211> 6
<212> PRT
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Ser Lys Leu Lys Ser Leu
1
<210> 206
<211> 8
<212> PRT
<213> Arabidopsis thaliana
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<221> VARIANT
<222> (1)...(8)
<223> Xaa at position 3 is Arg or His; Xaa at position 7
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<400> 206
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<210> 207
<211> 6
<212> PRT
<213> Arabidopsis thaliana
<400> 207
Ser Lys Leu Lys Ser Leu
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<210> 208
<211> 7
<212> PRT
<213> Arabidopsis thaliana
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Lys Phe Ser Tyr Asp Asn Leu
                5
<210> 209
<211> 23
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<213> Arabidopsis Thalia
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<222> 2,3,5,6,8,9,11,12,14,16+9,21,22
<223> Xaa=any amino acid
<221> VARIANT
<222> 4,15,20,23
<223> Xaa=L or I or V
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                5
Xaa Xaa Xaa Xaa Xaa Xaa
           20
<210> 210
<211> 23
<212> PRT
<213> Yeast
<220>
<221> VARIANT
<222> 2,3,5,6,8,9,11,12,14,16,17,19,21,22
<223> Xaa= any amino acid
<221> VARIANT
<222> 4,20,23
<223> Xaa=L or I or V
<400> 210
Pro Xaa Xaa Xaa Xaa Leu Xaa Xaa Leu Xaa Leu Xaa
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Xaa Asn Xaa Xaa Xaa Xaa
           20
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<211> 12
<212> PRT
<213> Arabidopsis thaliana
<220>
<221> VARIANT
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<222> 2,3,5,6,8,9,11
<223> Xaa=any amino acid
<221> VARIANT
<222> 1
<223> Xaa=I or L or V
<221> VARIANT
<222> 10
<223> Xaa=I or L
<400> 211
Xaa Xaa Xaa Leu Xaa Xaa Leu Xaa Xaa Xaa Leu
                5
                                    10
<210> 212
<211> 7
<212> PRT
<213> Arabidopsis thaliana
<220>
<221> VARIANT
<222> 1
<223> Xaa=I or R
<221> VARIANT
<222> 2,5-7
<223> Xaa=any amino acid
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                5
 1
<210> 213
<211> 8
<212> PRT
<213> Arabidopsis thaliana
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                5
<210> 214
<211> 16
<212> PRT
<213> Arabidopsis thaliana
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